

AMENDMENTS TO SPECIFICATION

Please amend the paragraph at page 22, lines 12-22 as follows:

The NH₂-terminal amino acid sequence of recombinant HPI isolated and analyzed as described above is set out in residues 1 to 40 of SEQ ID NO: 2. Degenerate oligonucleotide primers designed from this sequence were used to obtain the cDNA sequence set out in SEQ ID NO: 1 using cloning procedures described above. The full amino acid sequence comprising residues 1 to 40 of the purified native protein and deduced residues 41 to 181 of the cDNA is set out in SEQ ID NO: 2. The predicted molecular weight of this recombinant polypeptide is 20,333 Da, and its theoretical isoelectric point, 4.76. The estimated molecular weight of native HPI is about 20,314 to 20,365 Da. The HPI sequence was compared to other GENBANK sequences (~~published at www.ncbi.nlm.nih.gov~~), and found to exhibit little or no homology to previously published sequences.